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RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/828,313DATE: 08/27/2001
TIME: 12:31:16Input Set : A:\16313032.app
Output Set: N:\CRF3\08272001\I828313.raw

ENTERED

p. 5

3 <110> APPLICANT: COSTA e SILVA, OSWALDO DA
4 BOHNERT, HANS J.
5 THIELEN, NOCHA VAN
6 CHEN, ROUYING
7 SARRIA-MILLAN, RODRIGO
9 <120> TITLE OF INVENTION: PROTEIN KINASE STRESS-RELATED PROTEINS AND METHODS OF
10 USE IN PLANTS
12 <130> FILE REFERENCE: 16313-0032
14 <140> CURRENT APPLICATION NUMBER: 09/828,313
15 <141> CURRENT FILING DATE: 2001-04-06
17 <150> PRIOR APPLICATION NUMBER: 60/196,001
18 <151> PRIOR FILING DATE: 2000-04-07
20 <160> NUMBER OF SEQ ID NOS: 128
22 <170> SOFTWARE: PatentIn Ver. 2.1
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25 <211> LENGTH: 695
26 <212> TYPE: DNA
27 <213> ORGANISM: Physcomitrella patens
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31 <222> LOCATION: (636)
32 <223> OTHER INFORMATION: a, t, c, g, other or unknown
34 <220> FEATURE:
35 <221> NAME/KEY: modified_base
36 <222> LOCATION: (680)
37 <223> OTHER INFORMATION: a, t, c, g, other or unknown
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42 catgtctgga acctataccg tgaagagtga cgtctacagc ttcgggtgtg taatgctgga 180
43 gctactcact gggcgcaagc cttagacag ctcaagacca cgatccgagc aatctttggt 240
44 acgatgggcc acacctcaat tgcacgacat cgacgccctt gcacgaatgg tggatccgtc 300
45 gttgaagggc atctaccctg ctaaatacact ctctcgggtt gctgatatag tcgccctttg 360
46 cgtccagccg gagcccgagt tccgaccccc gatgtctgaa gtggtgcagg cacttgtaag 420
47 gctgatgcag cgtgcgagtc tgagcaaacg cagatcggag tccgctgttg ggaattgagt 480
48 cgaacgagcc atctgagact tcacctttga gagtactgaa gcgcccacta gcctaatacgt 540
49 gcatcttttg ccatctcgtt tctgagtgga acacaaagct gggatatattc tttggtggtt 600
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W--> 51 tgcaaagcgt tcgcgtggtn tgcttgaact tcaaa 695
54 <210> SEQ ID NO: 2
55 <211> LENGTH: 512
56 <212> TYPE: DNA
57 <213> ORGANISM: Physcomitrella patens
59 <400> SEQUENCE: 2
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61 gacaatctgg cctggtgtta gccagcaccg tgattggcac gagtttcctc aatggagacc 120
62 acaagatctg tcccttgctg ttcccgagct cagcgcggtt ggcttagacc ttctcgccaa 180

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63 aatgttggtta ttcgagccct caaagagaat ctctgccaaa gccgccttga gccatactta 240
64 ttctcgctgat gttgataaga cagcaaccta aacacaacag aacaattcaa gagaaccagg 300
65 taacctctac ctgtccaaga cgaaggacat ctaactcttc agtcaaactt ggccaatcat 360
66 gctgattggg aattgaacca caggaacgag gtgggcaccg tggttcgctg tagcatacaa 420
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73 <212> TYPE: DNA

74 <213> ORGANISM: Physcomitrella patens

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77 <221> NAME/KEY: modified_base

78 <222> LOCATION: (608)

79 <223> OTHER INFORMATION: a, t, c, g, other or unknown

81 <400> SEQUENCE: 3

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84 tctccgagag gcgaggcgga gagtgcagga tacatggaga tggtaaagct gttgaaggat 180
85 tacggggctg agtcacacgc aggtgccccg aggggccacg ttgagagtct gattcagggt 240
86 gccctccgt tgccttctaa ccgcgactgg gagatcgctc cgtcggagat tgaacttgat 300
87 accagcgagc tcatcggaag aggtcctttt ggagagattc ggaaggcgct ttggcgcggc 360
88 acaccgctcg ctgtgaagac aatcagacct tctctgtcca acgacagaat ggtcatcaag 420
89 gacttccagc acgaggtgca attgctcgta aaggttcggc acccaaacat tgtgcagttc 480
90 ctcggggctg ttaccggtca aagacctctc atgttagtca ccgagtttct ggcagggggg 540
91 cgatttgcat cagttgctga ggagcaccct aaatttggtt cctgaccgca tcgtgaagta 600
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96 <211> LENGTH: 710

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98 <213> ORGANISM: Physcomitrella patens

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101 <221> NAME/KEY: modified_base

102 <222> LOCATION: (54)

103 <223> OTHER INFORMATION: a, t, c, g, other or unknown

105 <400> SEQUENCE: 4

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107 gaagaaacgt aactccgtac ggctatttta ccaaattttc aagctcggtg tcccgccatg 120
108 atccaaatgg aagctcagtt tgcaacatga agtacattga acacacctac cgcccaccag 180
109 tcagaagcca ggccatgacc ttgtccttga atgatctcgg gtgctaagaa atcagccatg 240
110 ccacagactg tgaaagtgcg ctcacccgac atttgctttg caaaccgaaa atcaaccagc 300
111 tgaagtcgtc ctttccgac tatcataaga acatcgagg agatgccacg atatacaacg 360
112 ccattccttg gcagaagttc gacggctaat accacgttgg cgaccagaaa acgagctgag 420
113 ttctcgtcta aaggtgaccg aagtagaagt tctagaggcc cagctaacac acaattaaga 480
114 acgagtgcc caattgtcact gtcaataggg gtggccaaga gatcgggcac gaatggggaa 540
115 ggcctcagtt gcttgaaaag agttctctcc aataggactt ggccctccc accgagttc 600
116 tgaactttac gtctctggta ccttttcatg cttatgacgt catctgattt cttgcagagc 660
117 accacaccga catcacagca atcggttgaa tagacctggt gccgattcct 710
120 <210> SEQ ID NO: 5
121 <211> LENGTH: 1271

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122 <212> TYPE: DNA

123 <213> ORGANISM: Physcomitrella patens

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126 <221> NAME/KEY: modified_base

127 <222> LOCATION: (619)

128 <223> OTHER INFORMATION: a, t, c, g, other or unknown

130 <400> SEQUENCE: 5

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133 ataaagtcac ttctagtctg atccatacaa gctaccgaca caatgctaga agccttgatt 180
134 tacacactac aactagagaga gtctacaact cttttcctac actctgctta gttgcctcat 240
135 cctcaactcc ataaaccccc attcacaatc atgtaagact tgagagaggg aaacagtaag 300
136 caaccttggtg ctatttttagt accagagcag aggatgaacc actagtcctc ccaacgtaag 360
137 ccctaattcg ccgcaacaac ctacgacgg aactccgact tggtaaggg tggacaatat 420
138 gatacattcg aaggctgatt ttgcaaatgg gacgaagcag cgggaattctg gctgcgcact 480
139 gattgcagag agccattctg ggggagttga gtatacacag tccagtcgta cacatggctg 540
140 agctggaatt ttttctgaat gaaaagatca cggacaagc ttcggaggta cagtagtcag 600
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142 taccatcac aaagctcctc ccacagtggg ggtcatcttg attttgttgt gcatgtactc 720
143 gtgttgcttc tggtcagtga gggcggtgcc cgccttccc ttgccatggc aaattgcctc 780
144 ttagaaagta cataagaatg taaccaagt gattctatgt catctcttct actgtgctcg 840
145 attcctctgt gctgattcct actagcgtac cgtgccgtcc ctgtgaagct cttcctatct 900
146 cggtaaggga tatgccttcg tgttgccggg tccatgtact ctttgccaa gccaaaatct 960
147 ataataaaca cttgggttcc ttgccgaccg cagcccatga ggaagttatc cggcttcagg 1020
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149 agcataatca cggctctcaa cgaaaacctt agccacaca ccttaaagag gtgcaacagg 1140
150 ttcggcccca ataggtctag caccatcaca ttgtagtctt ctgctgcttt tccgaacct 1200
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152 aactctcgtg c 1271

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155 <210> SEQ ID NO: 6

156 <211> LENGTH: 1910

157 <212> TYPE: DNA

158 <213> ORGANISM: Physcomitrella patens

160 <400> SEQUENCE: 6

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162 aacaagctaa tggctgcaag cagtgaacat actaccaagg gagagattct cactccctag 120
163 acttcatect cgtacgttac ttggcaagga ttatggttta gtgataaaaa gcttcacaag 180
164 ccggcaagca tgctggttgc ttctgctgca atctaattgat tatttcctta ggaatcgtat 240
165 ggcagagagc taccacacaa agcactgaca atgggtttgat ggtaacaaga tagagatcca 300
166 ttcattccta agtatgagag acctgtagtc ttagcaccat tgtaggacag aaccaccgtt 360
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168 ttgatagaag acagagcccg gctatataac cacaagccct gcgcctacct tataacggct 480
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172 ctgttcgaga ggaatctacg acttctgctg atgccctga gggcctgctg ctagatagga 720
173 cagctcgctt ggaggaagaa cccccccgag ttgcatacga agatgtatgc atgcgctctg 780
174 gttctgacac aacagcaaga gcagaatcct tagcagattc atcaagtcca ggacttttgt 840
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177 ttccaagtgc accactggtt tgccctgcag aagcaccag tcgagttgaa ctgccaccgg 1020
178 aaatttgtga ctgctggtac ttcagaattg tccagtcaaa aacgtagtca aattgaaaac 1080
179 ctgtaaaaact atttccagtt taggcaaaca gaagtggcac tgtaataaac tgaaaatcat 1140
180 caaacattca caaactatct gttcggttgat agagcatagt aaagtctgcg cttaggatca 1200
181 agtcttgata cattacaatg cccaagcaag agtgaaacct acaaaagtta cagttttcat 1260
182 accctcacga ataaagaggt cacggaagat tcttttcaaa tatgcatagt cgggtttgtc 1320
183 atcaaaacgc aaggaccggc agtagtgga gtacgctcgt gcgaattctg aaggataatt 1380
184 tttaaaaagg acctcaatgg gcgtggacat ttgttttctc actgatcttc tcgtacttct 1440
185 gcttcttggt tcccgttttc agtccttgcc catggaagac tgcctctcag gaagtacatg 1500
186 agcacatatc caagagattc caaatcatct cgtctgcttt gctcaatacc aagatgagtg 1560
187 ttgatgcttg cataccgagc agtccctgtc agatttttgt tctccctgta gggaaatagc 1620
188 tgatgcgtgg aagggtcgcg gtacttcttg gcaagaccaa aatcaataat gtagacctgg 1680
189 tttgctcgcc taccaagccc cattagaaaa ttatcaggct tgatgtctct atgaagaaag 1740
190 ctttttcgcat gcacatactc cactctgttg atcagctggt cagcaagcat gagaacagtc 1800
191 ttttaaagaga acttccggct gcagaagttg aaaaggctct cgagacttgg ccccaacaga 1860
192 tccagaacca agacattgta gtctccttct atcccgaaac atcctcgtgc 1910

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195 <210> SEQ ID NO: 7

196 <211> LENGTH: 720

197 <212> TYPE: DNA

198 <213> ORGANISM: Physcomitrella patens

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201 <221> NAME/KEY: modified_base

202 <222> LOCATION: (58)

203 <223> OTHER INFORMATION: a, t, c, g, other or unknown

205 <220> FEATURE:

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207 <222> LOCATION: (612)

208 <223> OTHER INFORMATION: a, t, c, g, other or unknown

210 <400> SEQUENCE: 7

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213 gaaaaaaaag atcaaaagag agatcaagat tctgcaaaac ctttgtggag ggccaacat 180
214 tgtgaagctt ctggacattg tccgtgatca gcaatcgaag acaccagcc taatttttga 240
215 gtatgtgaac aatactgatt tcaaagtgtc ctacccact cttacagact ttgatatccg 300
216 atactacatt catgagctgc tcaaggcttt ggactattgc cattctcaag ggattatgca 360
217 cagggatgtg aagccacaca acgtgatgat tgaccatgag cagcggaagc ttaggcttat 420
218 tgactgggga cttgccgaat tctatcatcc tggcaaagag tataatgtgc gtgttgccctc 480
219 taggtacttc aagggtcctg agctgctggt tgatcttcaa gattatgatt actctctcga 540
220 catgtggagc tctggggtgc atgtttgccg gcatgatatt tcggaaggag ccattctttt 600
W--> 221 atgggcatga caattcatga tcaacttggg gaagatcgct aagggtgttg gaacttgatg 660
222 aattgaattc ctatctaaca aataccgcta agtggacccc attggagcac ctggtggggg 720

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226 <211> LENGTH: 953

227 <212> TYPE: DNA

228 <213> ORGANISM: Physcomitrella patens

230 <400> SEQUENCE: 8

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232 taagttgcaa aaccatacaa cgccgggata gcgttgtagc cacctaaaga attgagagta 120

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233 ggccttataa cttgagatga agtgtgaagt ggtactgcac catatcatca ggacctaagc 180
234 tgcaatccag agcctccctc caaatgagat ccctgatagg ctccctccgag atagagggct 240
235 cctcgaagcc aaactcgaag ggagataccg agccaggctc atcggtgatg tcatgaagtg 300
236 aagcttaaat aaggggtgcgc caaggcagct tccactgtga ttcttttcgc tggatcaaag 360
237 accagcatct tttcaacaag atcaagagca gaacgattaa tgccctctgaa cttctgggtt 420
238 aaggggaatag gcgactgtcg aggcagggtgc ttgatatacc gcctagcatt gtcgcttctc 480
239 aaaaacccaa gatccctatc ttcaggagtt ccgatgagtt ctgtaattag gcggagctga 540
240 tgcacatagt ctctcccagg gaacaacgca gatcggttaa gcaactccat gaagatgcac 600
241 cccacagacc aaatgtcaat agctgcagtg tatgctgaac aattcaggag cagctctgga 660
242 gctctgtacc acctcgttac aacatactca gtcatgaaat ccgtttcaga gagagtgcgt 720
243 gccaaagccaa aatctgcgat tttcaaatacg caattggcat tgacgagaag gttggtgggc 780
244 ttcaagtccc ggtgcaagac gtteggccgaa tggatgtact tcaagccccg caagatttga 840
245 tacagaaaat actgacagtg gtcttctgtg agagcttgat ttgaacgaat gatctggtgt 900
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250 <211> LENGTH: 683

251 <212> TYPE: DNA

252 <213> ORGANISM: Physcomitrella patens

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255 <221> NAME/KEY: modified_base

256 <222> LOCATION: (603)

257 <223> OTHER INFORMATION: a, t, c, g, other or unknown

259 <220> FEATURE:

260 <221> NAME/KEY: modified_base

261 <222> LOCATION: (610)

262 <223> OTHER INFORMATION: a, t, c, g, other or unknown

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267 taatctttcc ggagaggcct tatcgccgac ggcaactgcc gaagacgaga ttactcagat 180
268 gatactaaaa agtgccgcaa ggtccgaatt aggaatgtat gtttcgaaga gacaggaatt 240
269 ctatcttcga agagcgcgga ggcggcgtaa gtttgcggtg aagccgggtt tgcagagcat 300
270 ctccgagatg aagcctgtca tgggaattcca cactccgatg gcttaccggg atagtgggtc 360
271 tccgccgaag aacgcctcta ccccatcctt acctggcccc aagaacattt caccgccacg 420
272 acaagtgagt gtcccgcaaa ggagcagtc tccgccgaag aacgtctcac cacctcccca 480
273 gcccggcatt ttgtagcgcg gactgcatc gaagtattct gctgcatctc agcaagttca 540
274 acgaaatcga gggcaacgcg aaatctcttt tatatggcgt agtttgtgtc tccgactgga 600
275 ctccatctta tcccatcga gataactgat tcggtggata atttctccaa attttggcta 660
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281 <212> TYPE: DNA

282 <213> ORGANISM: Physcomitrella patens

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285 <221> NAME/KEY: modified_base

286 <222> LOCATION: (923)

287 <223> OTHER INFORMATION: a, t, c, g, other or unknown

289 <220> FEATURE:

290 <221> NAME/KEY: modified_base

Use of n and / or Xaa has been detected in the Sequence Listing. Review the Sequence Listing to ensure a corresponding explanation is present in the <220> to <223> fields of each sequence using n or Xaa.

VERIFICATION SUMMARY

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L:51 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:92 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:106 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:141 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:211 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:221 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:276 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:310 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10
L:314 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10
L:387 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13